

SEQUENCE LISTING

<110> Baum, Peter Robert
Fanslow III, William C.

<120> Molecules Designated LDCAM

<130> 2873-US

<140> to be assigned--

<141> 2001-02-06

<140> PCT/US99/17905

<141> 1999-08-05

<150> 60/095,672

<151> 1998-08-07

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<170> PatentIn Ver. 2.0

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Asp Ser Gly Thr Tyr Gly Cys Thr Ala Thr Ser Asn Met Gly Ser Tyr
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 Lys Ala Tyr Tyr Thr Leu Asn Val Asn Asp Pro Ser Pro Val Pro Ser
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 tcc tcc agc acc tac cac gcc atc atc ggt ggg atc gtg gct ttc att 1278
 Ser Ser Ser Thr Tyr His Ala Ile Ile Gly Gly Ile Val Ala Phe Ile
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 gtc ttc ctg ctg ctc atc atg ctc atc ttc ctt ggc cac tac ttg atc 1326
 Val Phe Leu Leu Leu Ile Met Leu Ile Phe Leu Gly His Tyr Leu Ile
 375 380 385 390
 cgg cac aaa gga acc tac ctg aca cat gag gca aaa ggc tcc gac gat 1374
 Arg His Lys Gly Thr Tyr Leu Thr His Glu Ala Lys Gly Ser Asp Asp
 395 400 405
 gct cca gac gcg gac acg gcc atc atc aat gca gaa ggc ggg cag tca 1422
 Ala Pro Asp Ala Asp Thr Ala Ile Ile Asn Ala Glu Gly Gly Gln Ser
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 Gly Gly Asp Asp Lys Lys Glu Tyr Phe Ile
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 <213> Homo sapien

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 Ile Ser Ser Thr Val Trp Ser Ser Pro Asp Met Leu Ala Ser Gln Asp
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 Ser Gln Pro Trp Thr Ser Asp Glu Thr Val Val Ala Gly Gly Thr Val
 65 70 75 80
 Val Leu Lys Cys Gln Val Lys Asp His Glu Asp Ser Ser Leu Gln Trp
 85 90 95
 Ser Asn Pro Ala Gln Gln Thr Leu Tyr Phe Gly Glu Lys Arg Ala Leu
 100 105 110

Arg Asp Asn Arg Ile Gln Leu Val Thr Ser Thr Pro His Glu Leu Ser
 115 120 125
 Ile Ser Ile Ser Asn Val Ala Leu Ala Asp Glu Gly Glu Tyr Thr Cys
 130 135 140
 Ser Ile Phe Thr Met Pro Val Arg Thr Ala Lys Ser Leu Val Thr Val
 145 150 155 160
 Leu Gly Ile Pro Gln Lys Pro Ile Ile Thr Gly Tyr Lys Ser Ser Leu
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 Arg Glu Lys Asp Thr Ala Thr Leu Asn Cys Gln Ser Ser Gly Ser Lys
 180 185 190
 Pro Ala Ala Arg Leu Thr Trp Arg Lys Gly Asp Gln Glu Leu His Gly
 195 200 205
 Glu Pro Thr Arg Ile Gln Glu Asp Pro Asn Gly Lys Thr Phe Thr Val
 210 215 220
 Ser Ser Ser Val Thr Phe Gln Val Thr Arg Glu Asp Asp Gly Ala Ser
 225 230 235 240
 Ile Val Cys Ser Val Asn His Glu Ser Leu Lys Gly Ala Asp Arg Ser
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 Thr Ser Gln Arg Ile Glu Val Leu Tyr Thr Pro Thr Ala Met Ile Arg
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 Pro Asp Pro Pro His Pro Arg Glu Gly Gln Lys Leu Leu Leu His Cys
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 Gly Ile Val Ala Phe Ile Val Phe Leu Leu Leu Ile Met Leu Ile Phe
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 Leu Gly His Tyr Leu Ile Arg His Lys Gly Thr Tyr Leu Thr His Glu
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tcg ctc ctg ctc ctg ctc ctg ctc ttc gcc tgc tgc tgg gcg ccc ggc 222
Ser Leu Leu Leu Leu Leu Leu Leu Phe Ala Cys Cys Trp Ala Pro Gly
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Gly Ala Asn Leu Ser Gln Asp Asp Ser Gln Pro Trp Thr Ser Asp Glu
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aca gtg gtg gct ggt ggc acc gtg gtg ctc aag tgc caa gtg aaa gat 318
Thr Val Val Ala Gly Gly Thr Val Val Leu Lys Cys Gln Val Lys Asp
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cac gag gac tca tcc ctg caa tgg tct aac cct gct cag cag act ctc 366
His Glu Asp Ser Ser Leu Gln Trp Ser Asn Pro Ala Gln Gln Thr Leu
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Tyr Phe Gly Glu Lys Arg Ala Leu Arg Asp Asn Arg Ile Gln Leu Val
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acc tct acg ccc cac gag ctc agc atc agc atc agc aat gtg gcc ctg 462
Thr Ser Thr Pro His Glu Leu Ser Ile Ser Ile Ser Asn Val Ala Leu
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gca gac gag ggc gag tac acc tgc tca atc ttc act atg cct gtg cga 510
Ala Asp Glu Gly Glu Tyr Thr Cys Ser Ile Phe Thr Met Pro Val Arg
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atc act ggt tat aaa tct tca tta cgg gaa aaa gac aca gcc acc cta 606
Ile Thr Gly Tyr Lys Ser Ser Leu Arg Glu Lys Asp Thr Ala Thr Leu
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aac tgt cag tct tct ggg agc aag cct gca gcc cgg ctc acc tgg aga 654
Asn Cys Gln Ser Ser Gly Ser Lys Pro Ala Ala Arg Leu Thr Trp Arg
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Thr Arg Glu Asp Asp Gly Ala Ser Ile Val Cys Ser Val Asn His Glu
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tct cta aag gga gct gac aga tcc acc tct caa cgc att gaa gtt tta 846

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Tyr	Thr	Pro	Thr	Ala	Met	Ile	Arg	Pro	Asp	Pro	Pro	His	Pro	Arg	Glu		
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ggc	cag	aag	ctg	ttg	cta	cac	tgt	gag	ggg	cgc	ggc	aat	cca	gtc	ccc	942	
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Gln	Gln	Tyr	Leu	Trp	Glu	Lys	Glu	Gly	Ser	Val	Pro	Pro	Leu	Lys	Met		
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acc	cag	gag	agt	gcc	ctg	atc	ttc	cct	ttc	ctc	aac	aag	agt	gac	agt	1038	
Thr	Gln	Glu	Ser	Ala	Leu	Ile	Phe	Pro	Phe	Leu	Asn	Lys	Ser	Asp	Ser		
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ggc	acc	tac	ggc	tgc	aca	gcc	acc	agc	aac	atg	ggc	agc	tac	aag	gcc	1086	
Gly	Thr	Tyr	Gly	Cys	Thr	Ala	Thr	Ser	Asn	Met	Gly	Ser	Tyr	Lys	Ala		
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tac	tac	acc	ctc	aat	gtt	aat	gac	ccc	agt	ccg	gtg	ccc	tcc	tcc	tcc	1134	
Tyr	Tyr	Thr	Leu	Asn	Val	Asn	Asp	Pro	Ser	Pro	Val	Pro	Ser	Ser	Ser		
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agc	acc	tac	cac	gcc	atc	atc	ggt	ggg	atc	gtg	gct	ttc	att	gtc	ttc	1182	
Ser	Thr	Tyr	His	Ala	Ile	Ile	Gly	Gly	Ile	Val	Ala	Phe	Ile	Val	Phe		
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ctg	ctg	ctc	atc	atg	ctc	atc	ttc	ctt	ggc	cac	tac	ttg	atc	cgg	cac	1230	
Leu	Leu	Leu	Ile	Met	Leu	Ile	Phe	Leu	Gly	His	Tyr	Leu	Ile	Arg	His		
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Lys	Gly	Thr	Tyr	Leu	Thr	His	Glu	Ala	Lys	Gly	Ser	Asp	Asp	Ala	Pro		
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gac	gcg	gac	acg	gcc	atc	atc	aat	gca	gaa	ggc	ggg	cag	tca	gga	ggg	1326	
Asp	Ala	Asp	Thr	Ala	Ile	Ile	Asn	Ala	Glu	Gly	Gly	Gln	Ser	Gly	Gly		
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gac	gac	aag	aag	gaa	tat	ttc	atc	tagaggcgcc	tgcccacttc	ctgcgcccc						1380	
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 Lys Cys Gln Val Lys Asp His Glu Asp Ser Ser Leu Gln Trp Ser Asn
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 Pro Ala Gln Gln Thr Leu Tyr Phe Gly Glu Lys Arg Ala Leu Arg Asp
 65 70 75 80
 Asn Arg Ile Gln Leu Val Thr Ser Thr Pro His Glu Leu Ser Ile Ser
 85 90 95
 Ile Ser Asn Val Ala Leu Ala Asp Glu Gly Glu Tyr Thr Cys Ser Ile
 100 105 110
 Phe Thr Met Pro Val Arg Thr Ala Lys Ser Leu Val Thr Val Leu Gly
 115 120 125
 Ile Pro Gln Lys Pro Ile Ile Thr Gly Tyr Lys Ser Ser Leu Arg Glu
 130 135 140
 Lys Asp Thr Ala Thr Leu Asn Cys Gln Ser Ser Gly Ser Lys Pro Ala
 145 150 155 160
 Ala Arg Leu Thr Trp Arg Lys Gly Asp Gln Glu Leu His Gly Glu Pro
 165 170 175
 Thr Arg Ile Gln Glu Asp Pro Asn Gly Lys Thr Phe Thr Val Ser Ser
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 Ser Val Thr Phe Gln Val Thr Arg Glu Asp Asp Gly Ala Ser Ile Val
 195 200 205
 Cys Ser Val Asn His Glu Ser Leu Lys Gly Ala Asp Arg Ser Thr Ser
 210 215 220
 Gln Arg Ile Glu Val Leu Tyr Thr Pro Thr Ala Met Ile Arg Pro Asp
 225 230 235 240
 Pro Pro His Pro Arg Glu Gly Gln Lys Leu Leu Leu His Cys Glu Gly
 245 250 255
 Arg Gly Asn Pro Val Pro Gln Gln Tyr Leu Trp Glu Lys Glu Gly Ser
 260 265 270
 Val Pro Pro Leu Lys Met Thr Gln Glu Ser Ala Leu Ile Phe Pro Phe
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 Leu Asn Lys Ser Asp Ser Gly Thr Tyr Gly Cys Thr Ala Thr Ser Asn
 290 295 300
 Met Gly Ser Tyr Lys Ala Tyr Tyr Thr Leu Asn Val Asn Asp Pro Ser
 305 310 315 320
 Pro Val Pro Ser Ser Ser Ser Thr Tyr His Ala Ile Ile Gly Gly Ile
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 Val Ala Phe Ile Val Phe Leu Leu Leu Ile Met Leu Ile Phe Leu Gly
 340 345 350
 His Tyr Leu Ile Arg His Lys Gly Thr Tyr Leu Thr His Glu Ala Lys

